

SEQUENCE LISTING

<110> BML, INC.

<120> Method of Detecting Risk Factor for Onset of Diabetes

<130> PBM37

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<170> PatentIn Ver. 2.0

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cgg ctc tct agg aga gcc caa ctc tgt ctt ggc gtc agt atc ctg gtc 96
 Arg Leu Ser Arg Arg Ala Gln Leu Cys Leu Gly Val Ser Ile Leu Val
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ctg atc ctc gtc gtg ctc gcg gtg gtc gtc ccg agg tgg cgcc cag 144
 Leu Ile Leu Val Val Val Leu Ala Val Val Val Pro Arg Trp Arg Gln
 35 40 45

cag tgg agc ggt ccg ggc acc acc aag cgc ttt ccc gag acc gtc ctg 192
 Gln Trp Ser Gly Pro Gly Thr Thr Lys Arg Phe Pro Glu Thr Val Leu
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gcg cga tgc gtc aag tac act gaa att cat cct gag atg aga cat gta 240

Ala Arg Cys Val Lys Tyr Thr Glu Ile His Pro Glu Met Arg His Val			
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Asp Cys Gln Ser Val Trp Asp Ala Phe Lys Gly Ala Phe Ile Ser Lys			
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His Pro Cys Asn Ile Thr Glu Glu Asp Tyr Gln Pro Leu Met Lys Leu			
100	105	110	
gga act cag acc gta cct tgc aac aag att ctt ctt tgg agc aga ata			384
Gly Thr Gln Thr Val Pro Cys Asn Lys Ile Leu Leu Trp Ser Arg Ile			
115	120	125	
aaa gat ctg gcc cat cag ttc aca cag gtc cag cgg gac atg ttc acc			432
Lys Asp Leu Ala His Gln Phe Thr Gln Val Gln Arg Asp Met Phe Thr			
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Leu Glu Asp Thr Leu Leu Gly Tyr Leu Ala Asp Asp Leu Thr Trp Cys			
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ggt gaa ttc aac act tcc aaa ata aac tat caa tct tgc cca gac tgg			528
Gly Glu Phe Asn Thr Ser Lys Ile Asn Tyr Gln Ser Cys Pro Asp Trp			
165	170	175	
aga aag gac tgc agc aac aac cct gtt tca gta ttc tgg aaa acg gtt			576
Arg Lys Asp Cys Ser Asn Asn Pro Val Ser Val Phe Trp Lys Thr Val			
180	185	190	
tcc cgc agg ttt gca gaa gct gcc tgt gat gtg gtc cat gtg atg ctc			624
Ser Arg Arg Phe Ala Glu Ala Ala Cys Asp Val Val His Val Met Leu			
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aat gga tcc cgc agt aaa atc ttt gac aaa aac agc act ttt ggg agt			672
Asn Gly Ser Arg Ser Lys Ile Phe Asp Lys Asn Ser Thr Phe Gly Ser			
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gtg gaa gtc cat aat ttg caa cca gag aag gtt cag aca cta gag gcc			720
Val Glu Val His Asn Leu Gln Pro Glu Lys Val Gln Thr Leu Glu Ala			
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tgg gtg ata cat ggt gga aga gaa gat tcc aga gac tta tgc cag gat			768
Trp Val Ile His Gly Gly Arg Glu Asp Ser Arg Asp Leu Cys Gln Asp			
245	250	255	
ccc acc ata aaa gag ctg gaa tcg att ata agc aaa agg aat att caa			816
Pro Thr Ile Lys Glu Leu Glu Ser Ile Ile Ser Lys Arg Asn Ile Gln			

260

265

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Phe Ser Cys Lys Asn Ile Tyr Arg Pro Asp Lys Phe Leu Gln Cys Val
275 280 285

aaa aat cct gag gat tca tct tgc aca tct gag atc tgagccagtc 910
Lys Asn Pro Glu Asp Ser Ser Cys Thr Ser Glu Ile
290 295 300

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35 40 45

Gln Trp Ser Gly Pro Gly Thr Thr Lys Arg Phe Pro Glu Thr Val Leu
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Ala Arg Cys Val Lys Tyr Thr Glu Ile His Pro Glu Met Arg His Val
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Asp Cys Gln Ser Val Trp Asp Ala Phe Lys Gly Ala Phe Ile Ser Lys
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His Pro Cys Asn Ile Thr Glu Glu Asp Tyr Gln Pro Leu Met Lys Leu
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Gly Thr Gln Thr Val Pro Cys Asn Lys Ile Leu Leu Trp Ser Arg Ile
115 120 125

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Gly Glu Phe Asn Thr Ser Lys Ile Asn Tyr Gln Ser Cys Pro Asp Trp
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Arg Lys Asp Cys Ser Asn Asn Pro Val Ser Val Phe Trp Lys Thr Val
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Asn Gly Ser Arg Ser Lys Ile Phe Asp Lys Asn Ser Thr Phe Gly Ser
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Trp Val Ile His Gly Gly Arg Glu Asp Ser Arg Asp Leu Cys Gln Asp
245 250 255

Pro Thr Ile Lys Glu Leu Glu Ser Ile Ile Ser Lys Arg Asn Ile Gln
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